

FIG. 1

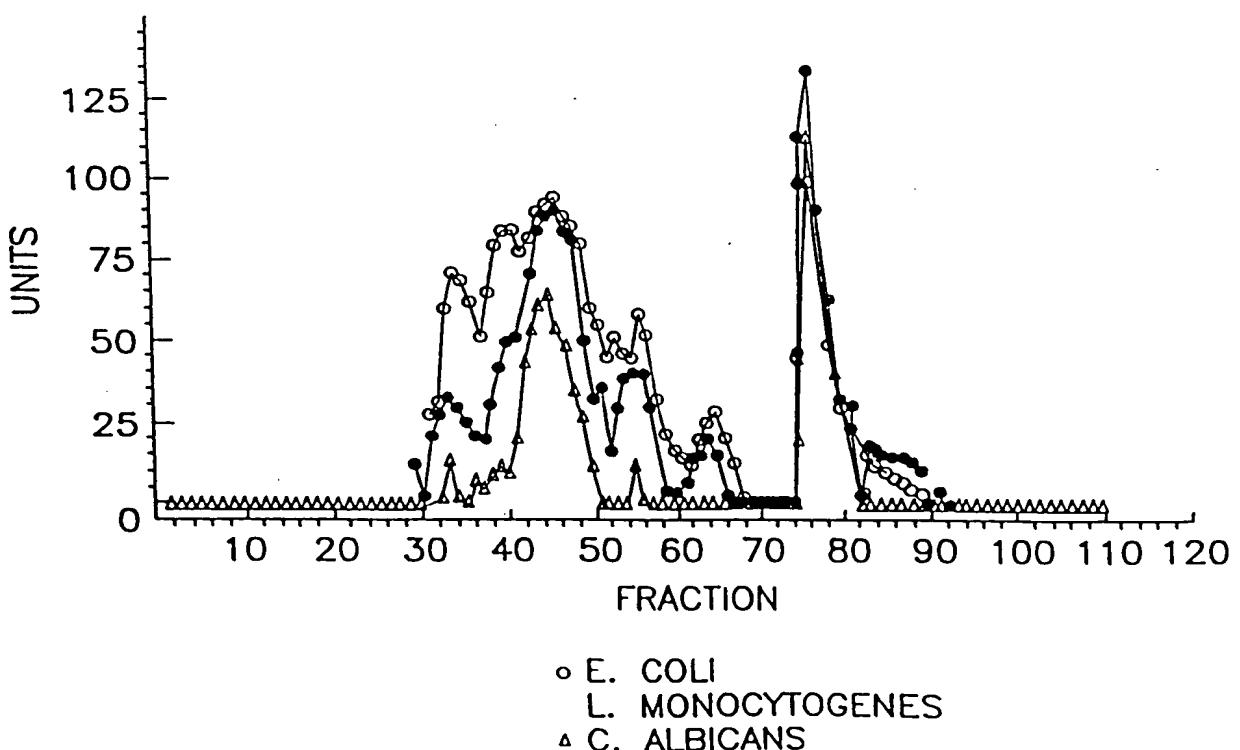


FIG. 2

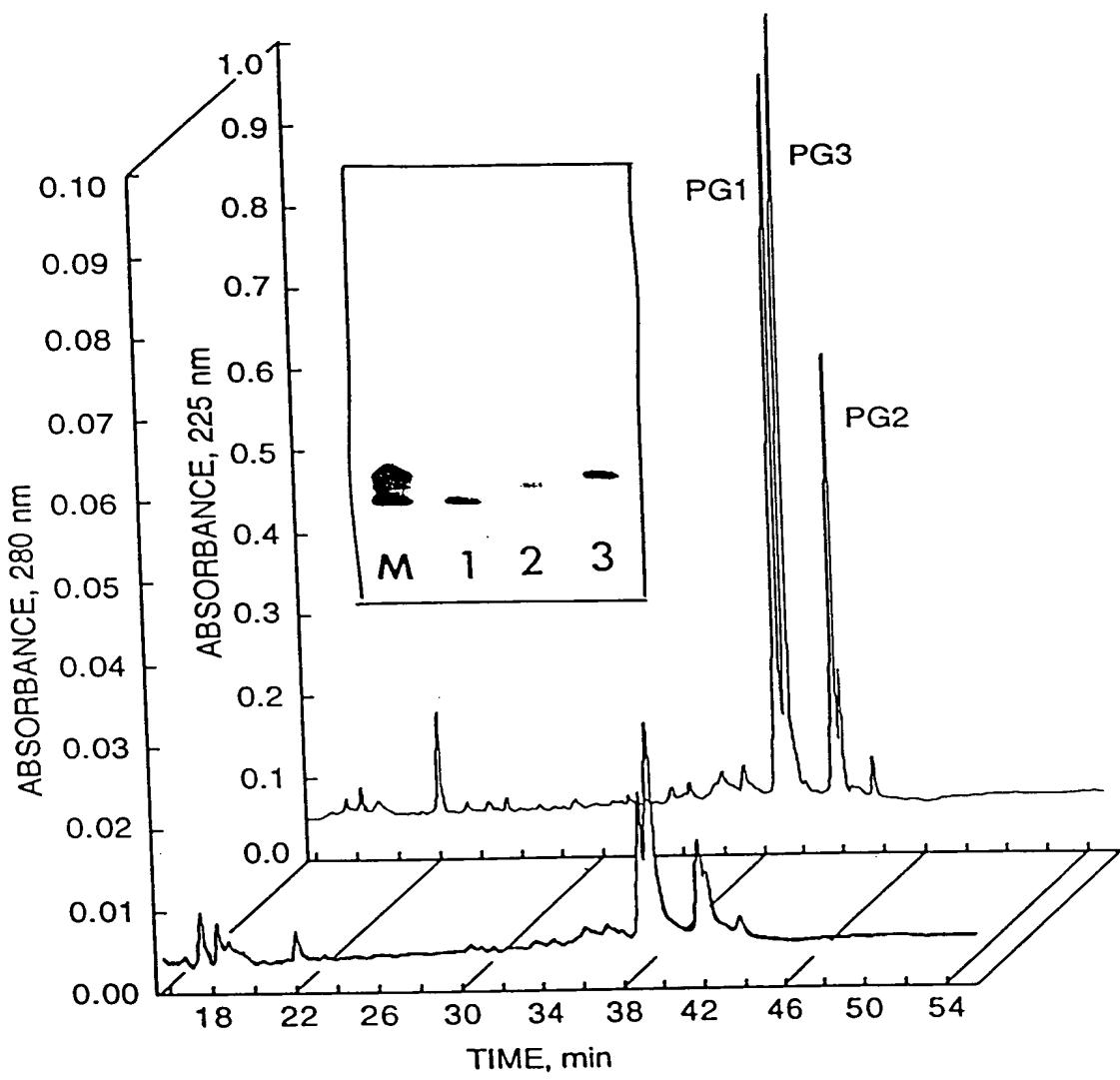


FIG. 3

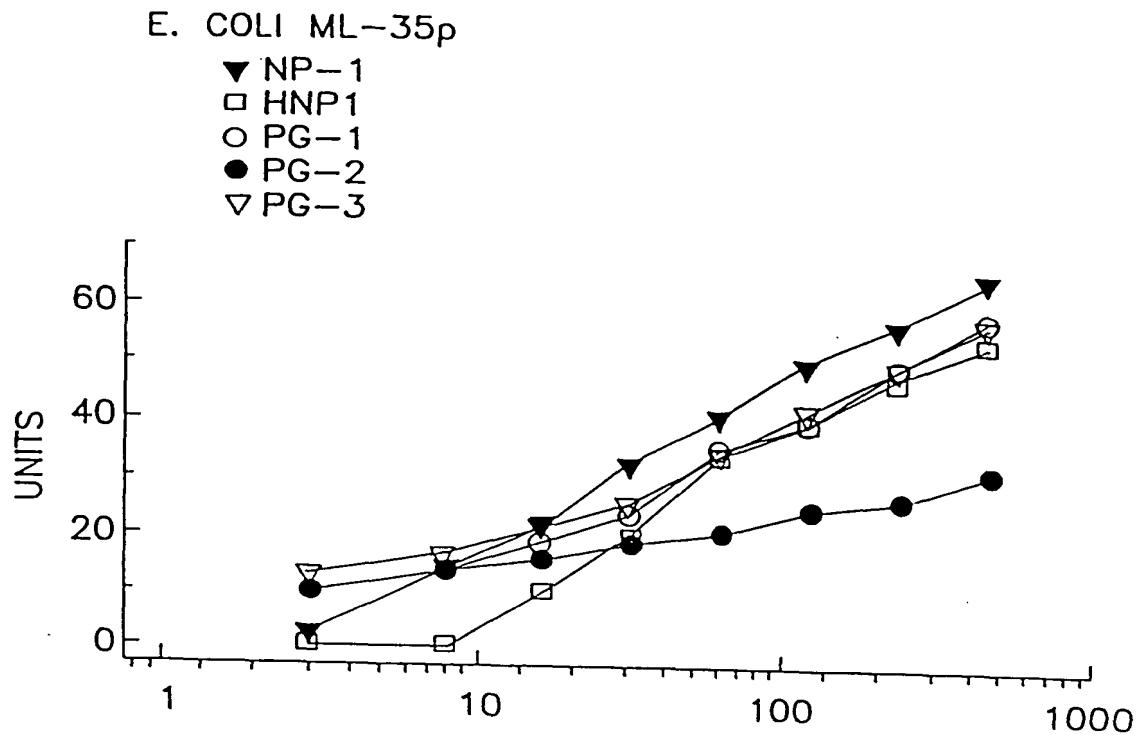


FIG. 4a

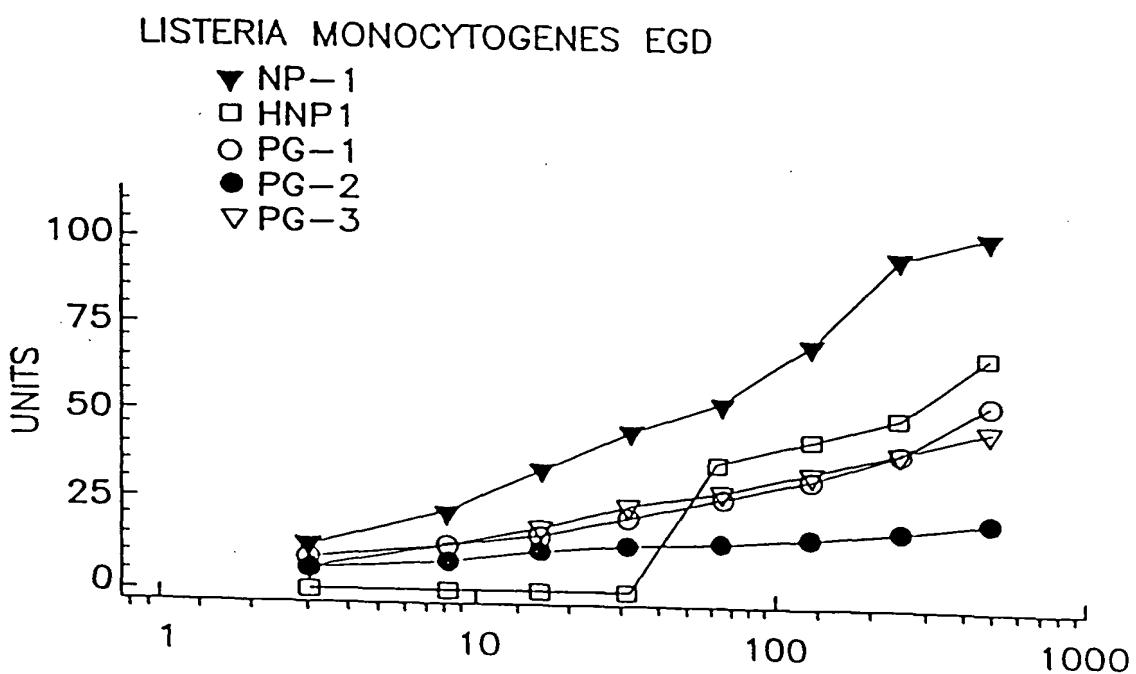


FIG. 4b

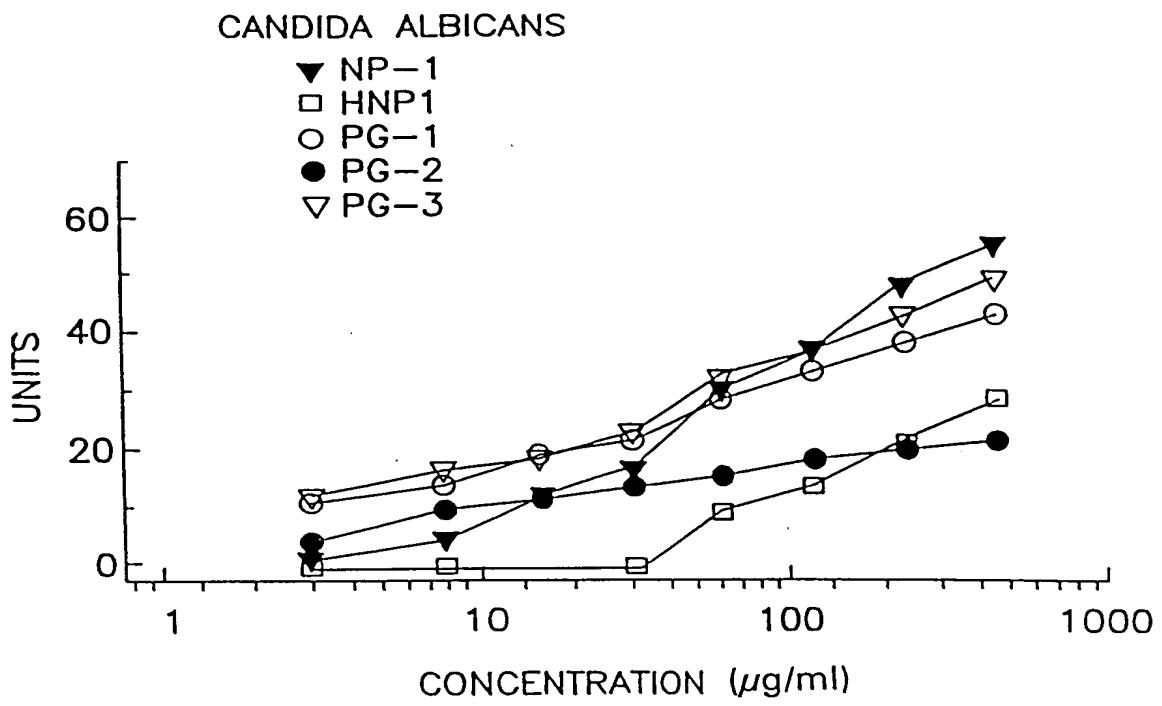


FIG. 4c

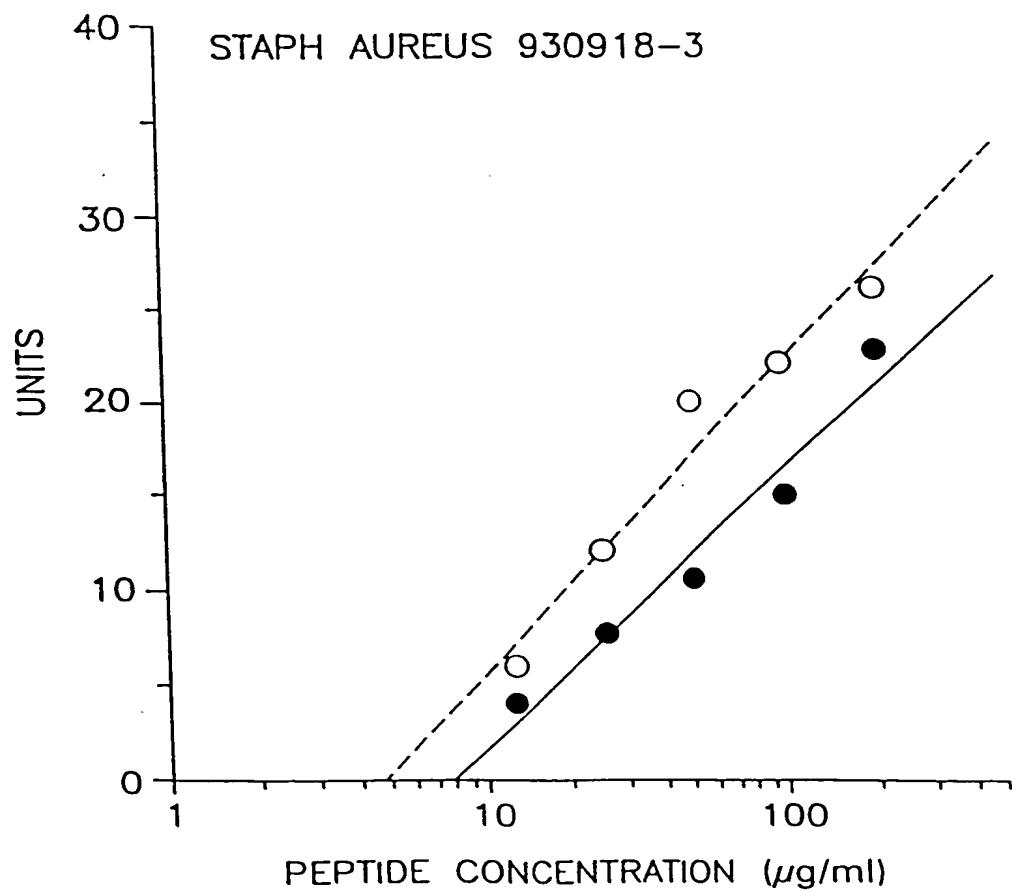


FIG. 4d

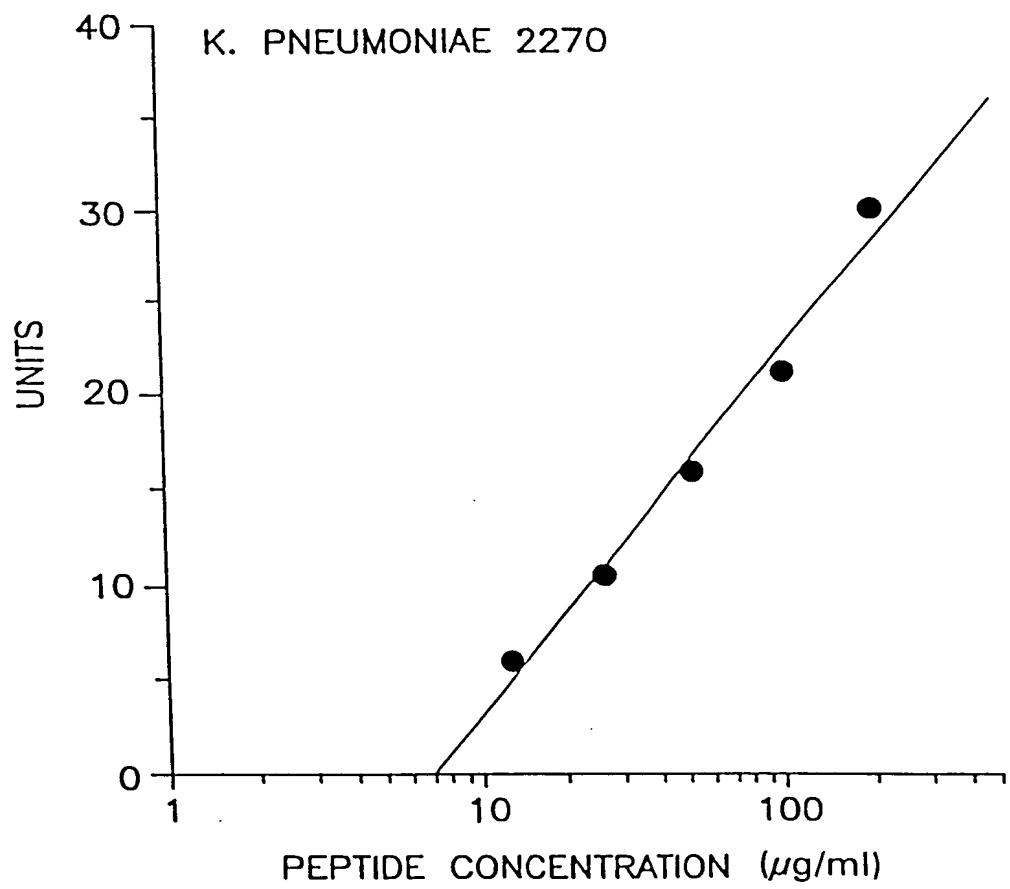


FIG. 4e

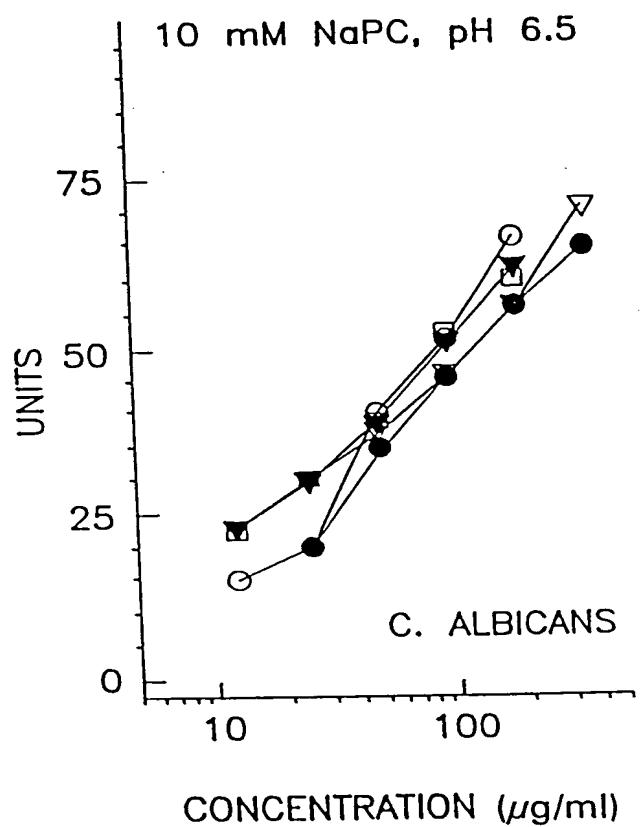


FIG. 5a-1

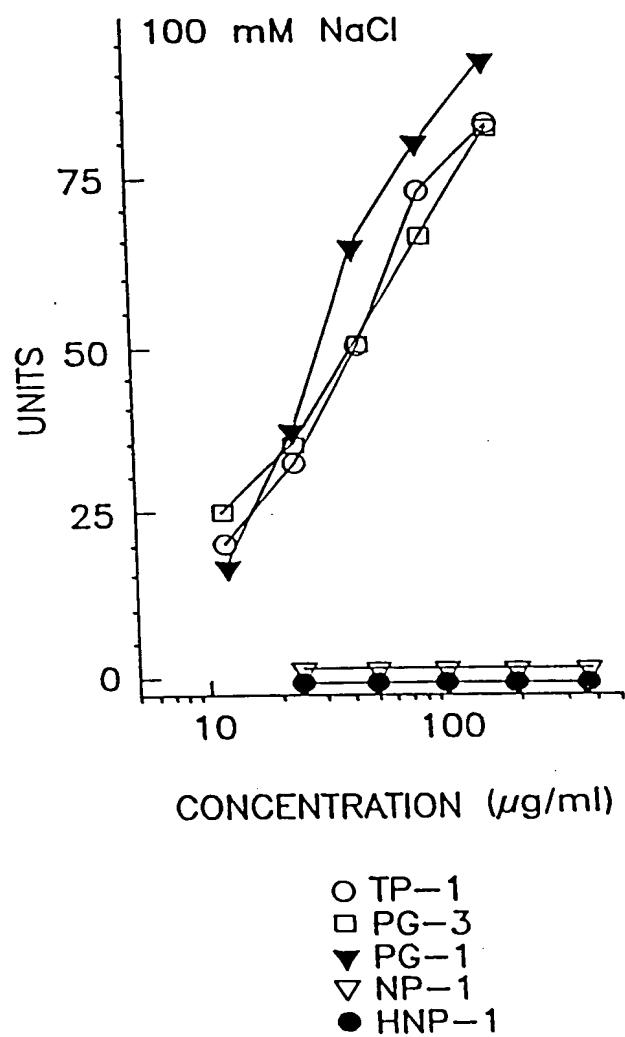


FIG. 5a-2

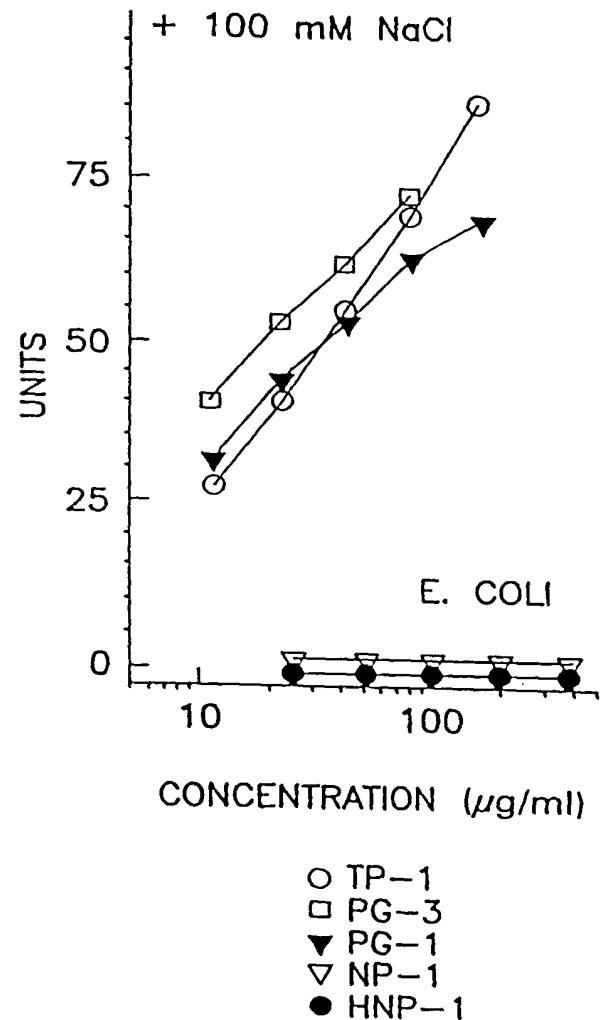
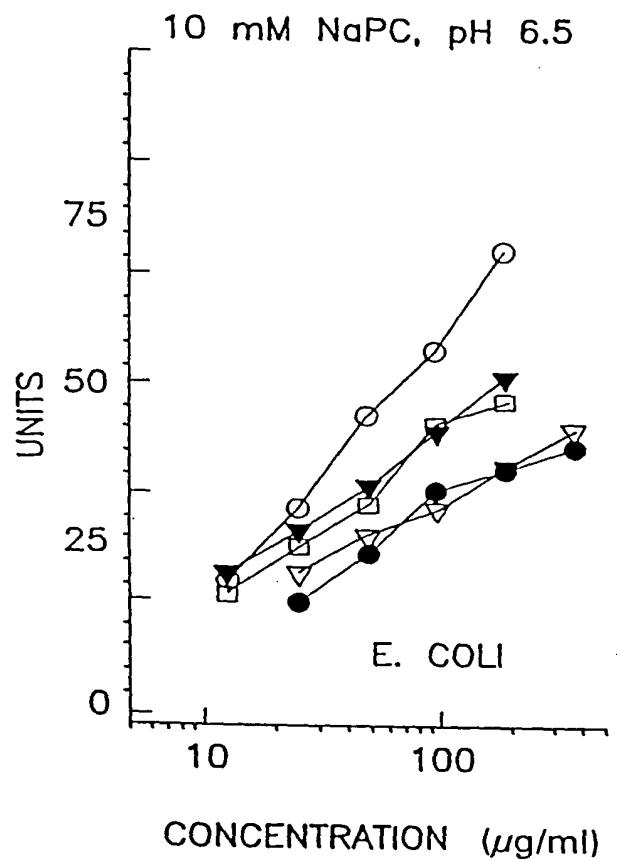


FIG. 5b-1

FIG. 5b-2

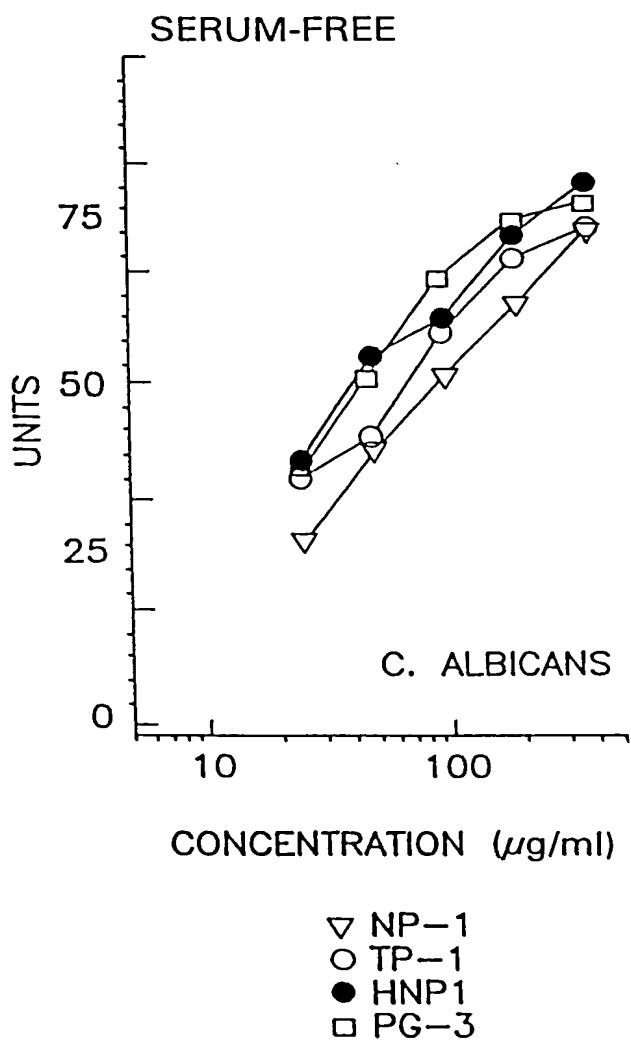


FIG. 5c-1

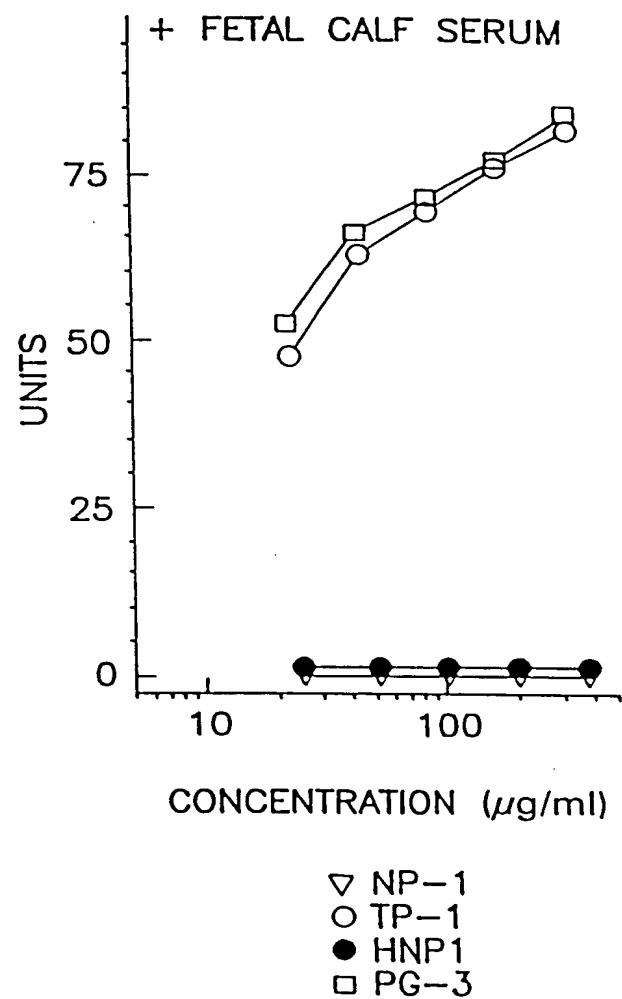


FIG. 5c-2

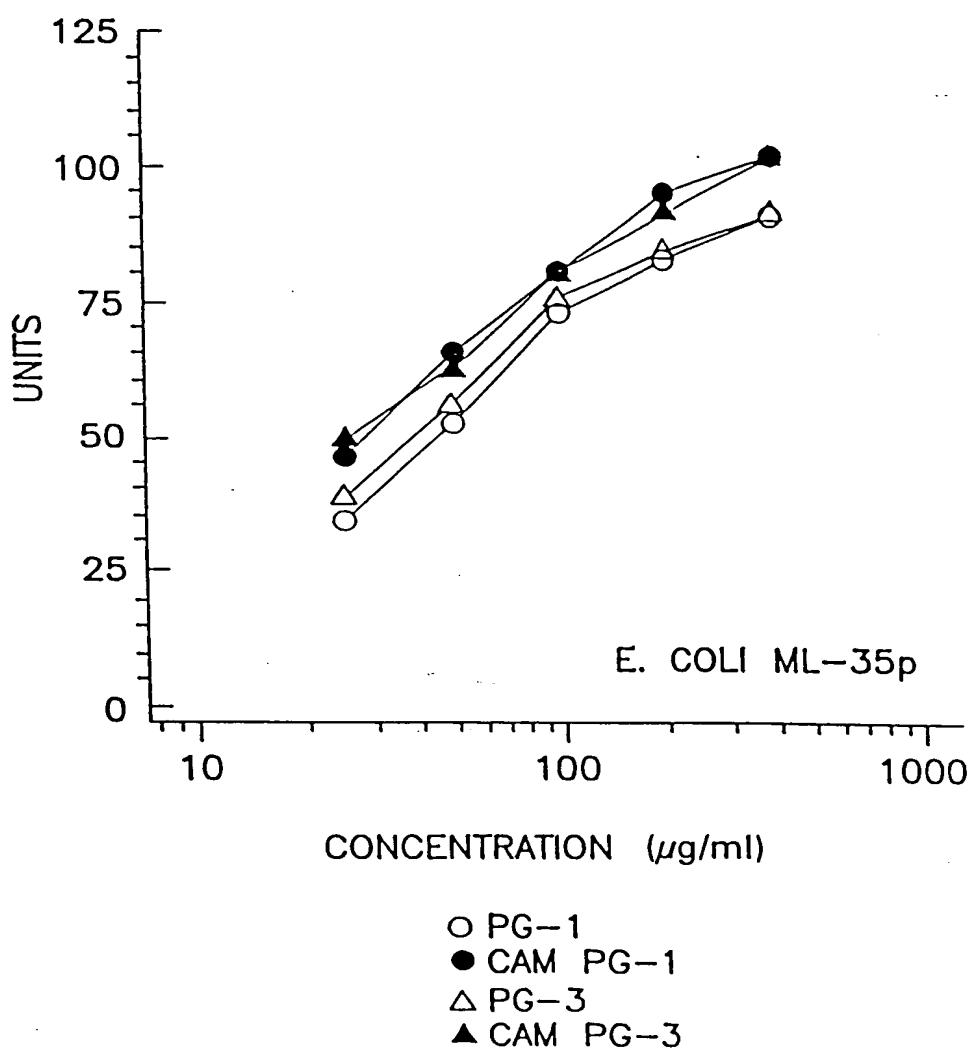


FIG. 6a

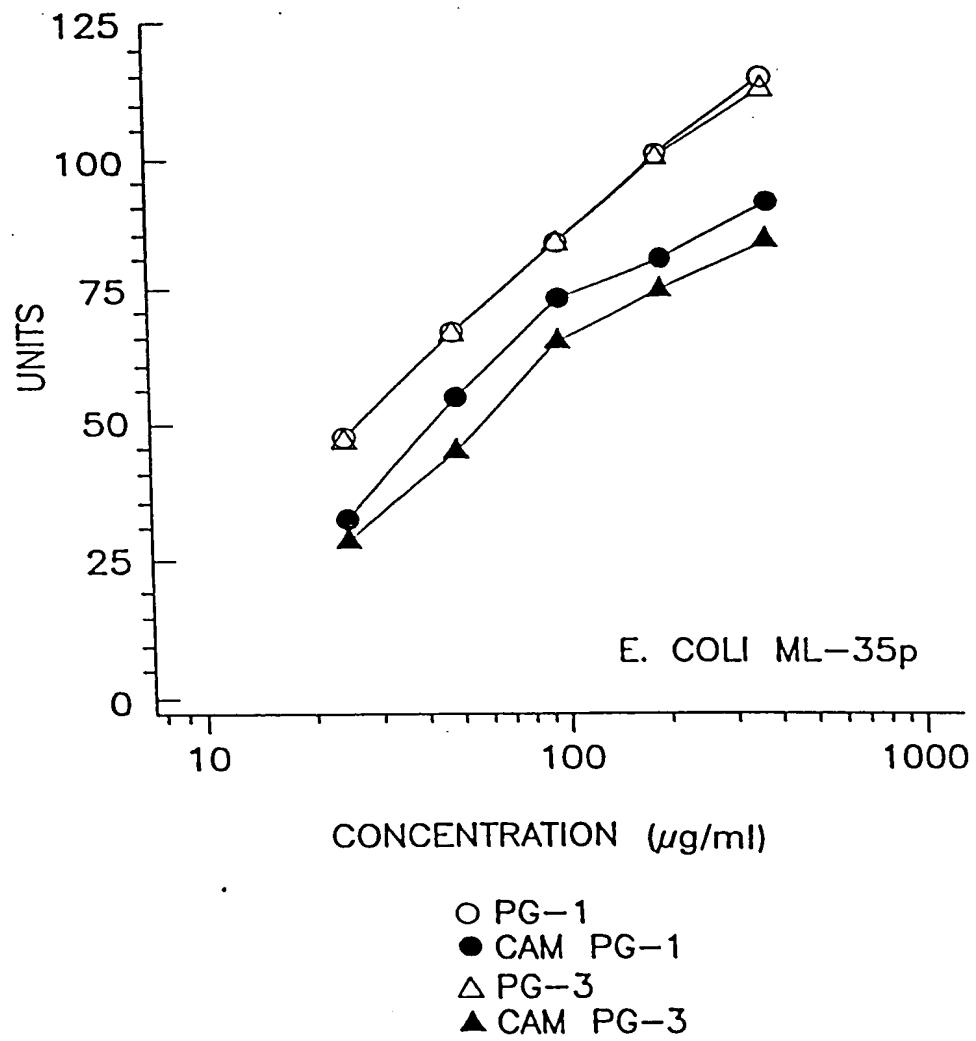


FIG. 6b

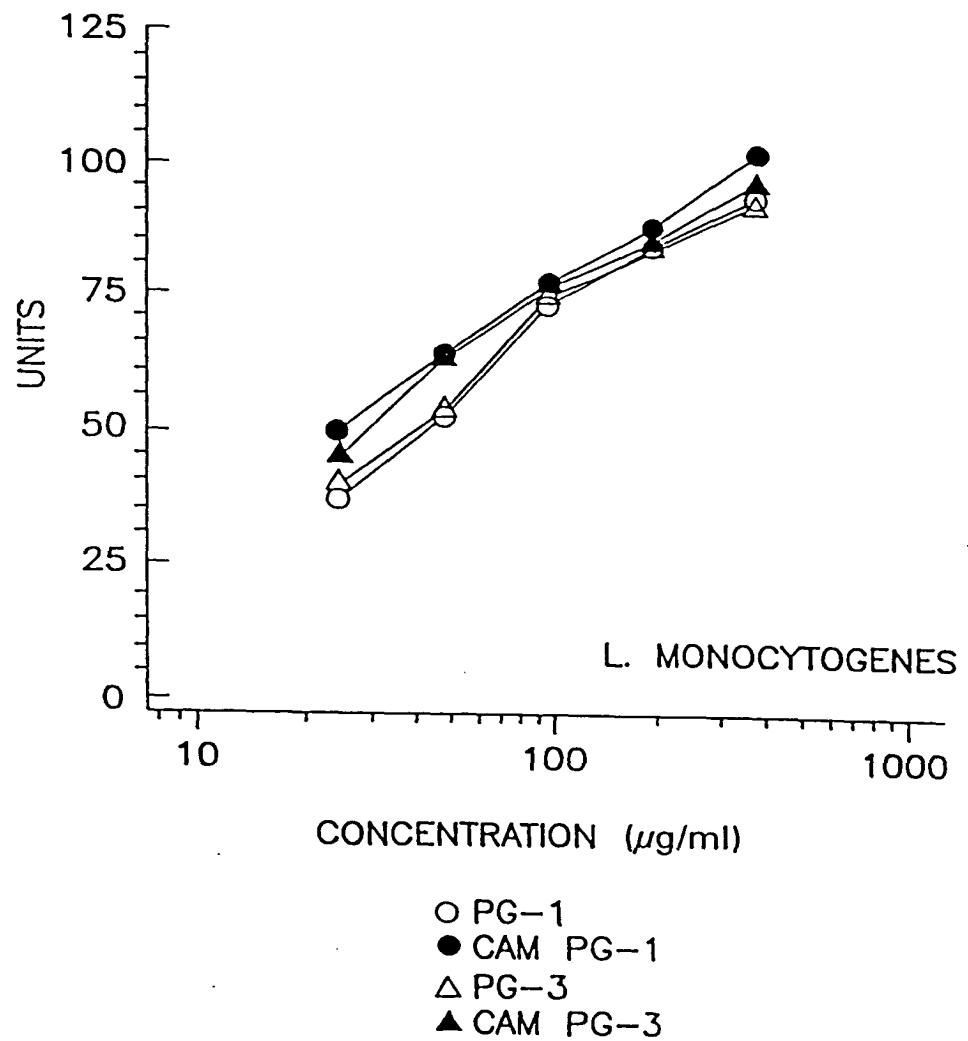


FIG. 6c

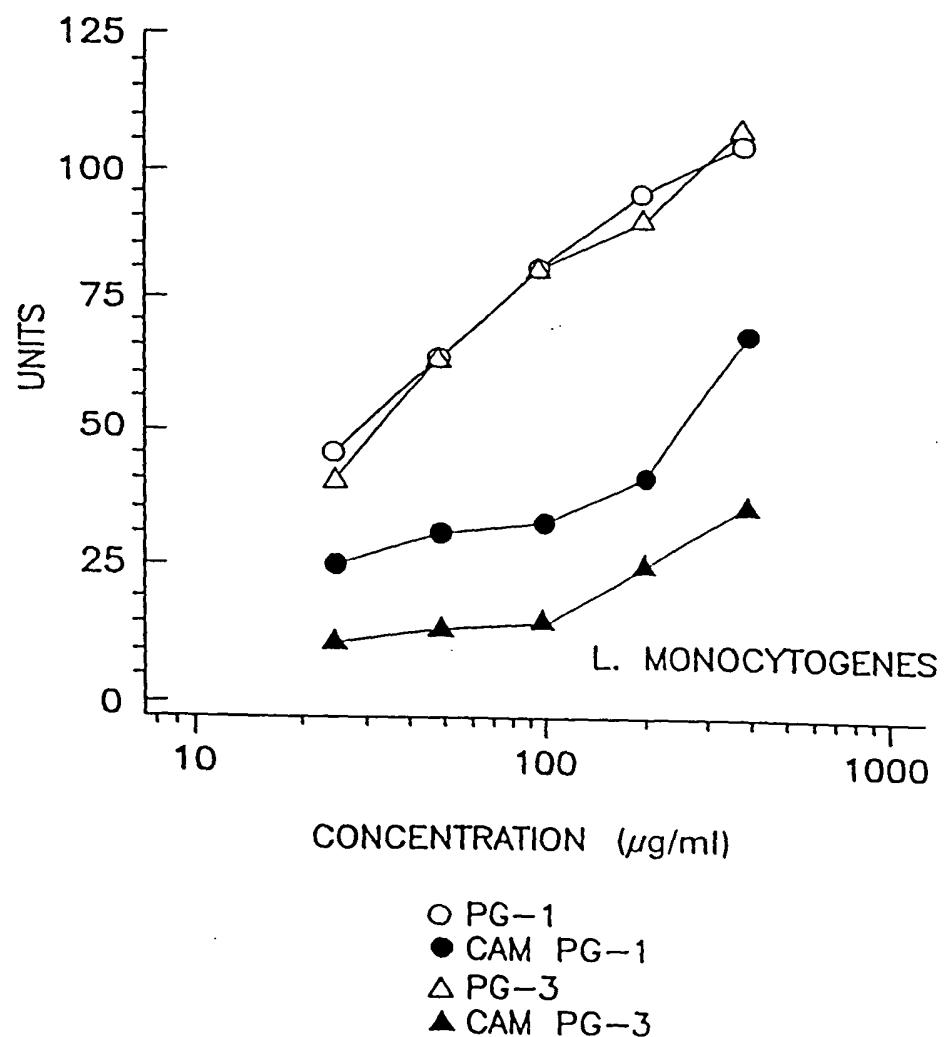


FIG. 6d

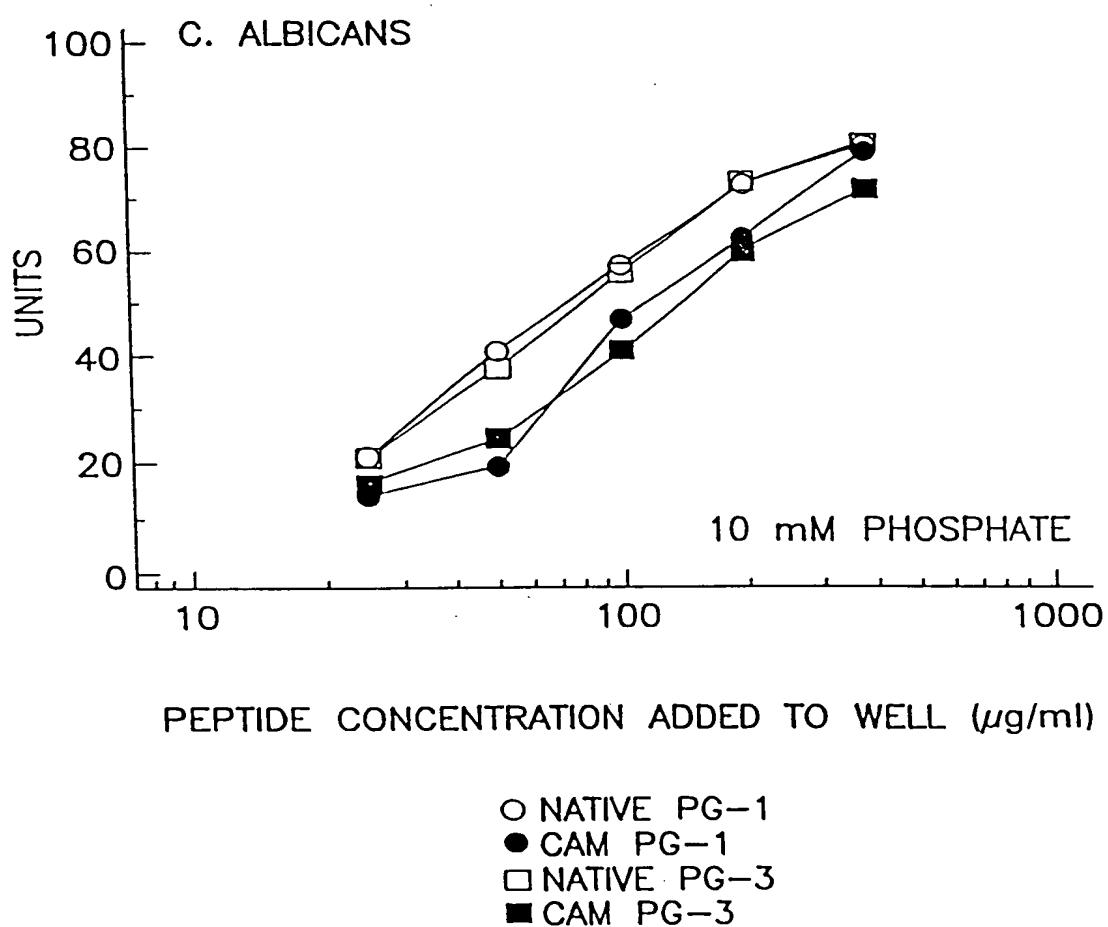


FIG. 6e

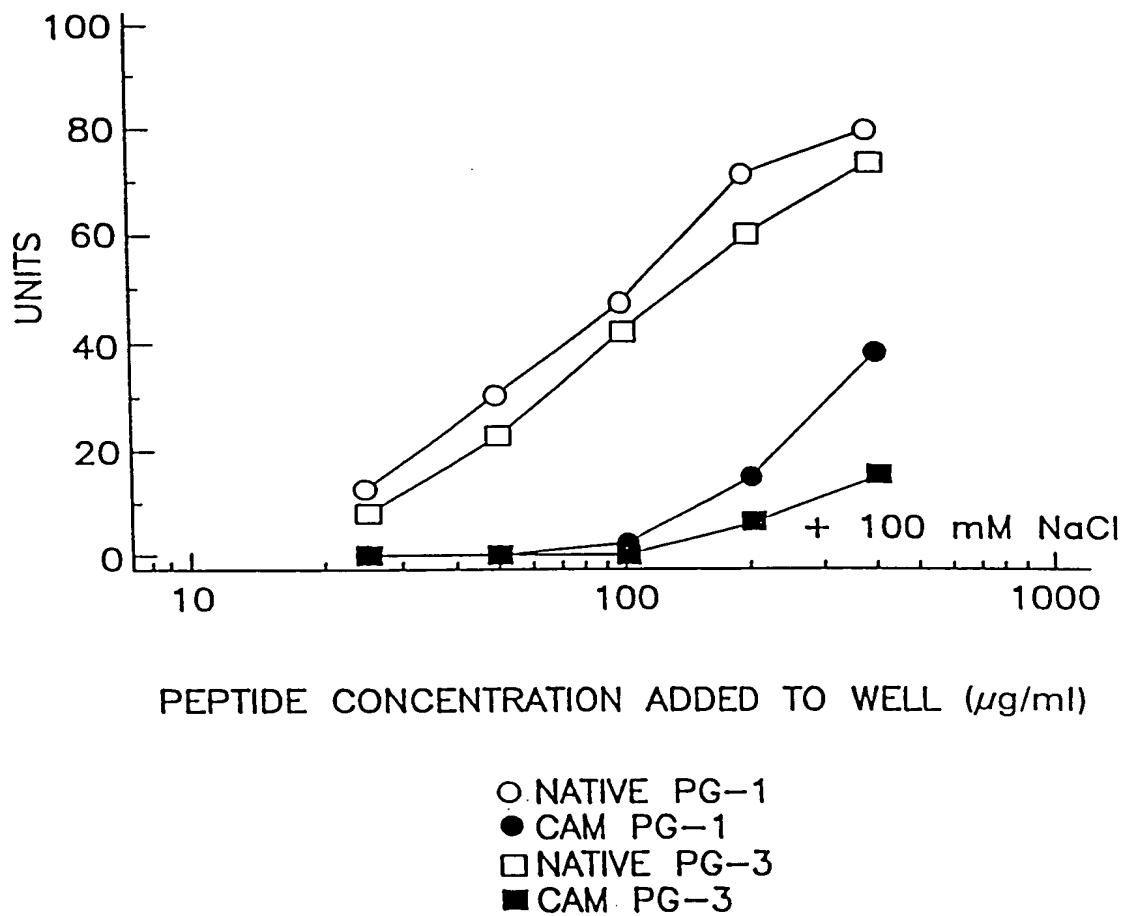


FIG. 6f

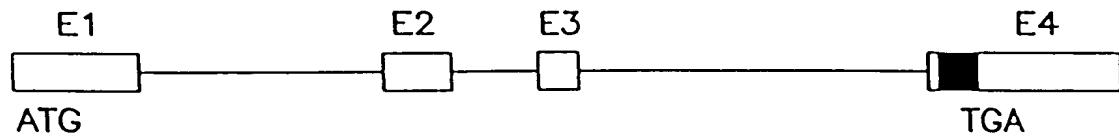


FIG. 9

10	20	30	40	50	
ATGGAGACCGAGAGAGCCAGCCTGTGCCCTGGGGCGCTGGTCACTGTGGCTTGCTGCTG					
MetGluThrGlnArgAlaSerLeuCysLeuGlyArgTrpSerLeuTrpLeuLeuLeuLeu					60 20
GCACTCGTGGTGCCCTCGGCCAGGCCAGGCCCTCAGCTACAGGGAGGCCGTGCTTCGT					120 40
AlaLeuValValProSerAlaSerAlaGlnAlaLeuSerTyrArgGluAlaValLeuArg					
GCTGTGGATCGCCTAACGAGCAGTCCTCGGAAGCTAATCTTACCGCCTGGAGCTG					180 60
AlaValAspArgLeuAsnGluGlnSerSerGluAlaAsnLeuTyrArgLeuLeuGluLeu					
GACCAGCCGCCAAGGCCGACGAGGACCCGGCACCCCGAAACCTGTGAGCTTCACGGTG					240 80
AspGlnProProLysAlaAspGluAspProGlyThrProLysProValSerPheThrVal					
AAGGGAGACTGTGTCCCAGGCCGACCCGGCAGCCCCGGAGCTGTGACTTCAAGGAG					300 100
LysGluThrValCysProArgProThrArgGlnProProGluLeuCysAspPheLysGlu					
AACGGGCGGGTGAAACAGTGTGTGGGGACAGTCACCTGGATCAGATCAAGGACCCGCTC					360 120
AsnGlyArgValLysGlnCysValGlyThrValThrLeuAspGlnIleLysAspProLeu					
^{G3}					^{G4}
GACATCACCTGCAATGAGGTTCAAGGTGTCAGGGAGGTGCCTGTGCTATTGTAGGCCT					420 140
AspIleThrCysAsnGluValGlnGlyValArgGlyGlyArgLeuCysTyrCysArgArg					
Gly ³					Gly ⁴
T ⁴ A ⁴ A ² T ⁴ T ²					
AGGTTCTGCGTCTGTGTCGGACGAGGATGACGGTTGCGACGGCAGGCTTCCCTCCCCA					480 149
ArgPheCysValCysValGlyArgGly---					
Trp ⁴ Ile ⁴ Phe ⁴ Ile ²	---2				
ATTTTCCCGGGGCCAGGTTCCGTCCCCAATTTTCCGCCTCACCTTCCGGCCCGCA					540
CCATTGGTCCACCAAGGTTCCCTGGTAGACGGTGAAGGATTGCAGGCAACTCACCCAG					600
AAGGCCTTCGGTACATTAAAATCCCAGCAAGGAGACCTAACGCATCTGCTTGCCAGGC					660
CCGCATCTGTCAAATAAATTCTTGTGAAACC					691

FIG. 7

ATGGAGACCCAGAGAGCCAGCCTGTGCCTGGGGCGCTGGTCACTGTGGCTCTGCTGCTG	60
M E T Q R A S L C L G R W S L W L L L	
^{G5}	
GCACTCGTGGTGCCCTCGGCCAGCGCCCAGGCCCTCAGCTACAGGGAGGCCGTGCTTCGT	120
A L V V P S A S A Q . A L . S . Y . R . E . A . V . L . R.	
^{G5}	
GCTGTGGATCGCCTAACGAGCAGTCCTCGGAAGCTAATCTTACCGCCTCTGGAGCTG	180
A V D R L N E Q . S . S . E . A . N . L . Y . R . L . L . E . L	
GACCAGCCGCCAAGGCCgtgagtcgggcaggggctcaggaggggctggggggcggggc	240
D Q P P K A	
tgtccccccacccggcccccgggctccctgtccctcccccgtcaggctgtccctcccgcc	300
aggaaggcacttgtccctctaagggggacccccttgccaggaaaccttcccagagctgg	360
gtgccctgcccgcgtgagagcttcccgccttagcctctggctgtggctcaggccctg	420
cacagcctgtgaggcaggagcgggctctgtccctcccccgtgcacccagcaccaagccc..	480
aggccaggctcccagcaggggctgcagaggctgtgtctaggtggggcggggaggggg	540
tgacagatccgagggggaagccctgagcccggatccatctcccaactttgatccttgacc	600
^{A5}	
agGACGAGGACCCGGGCACCCCCGAAACCTGTGAGCTCACGGTGAAAGGAGACTGTGTGTC	660
D E D P G T P K P V S F T V K E T V C	
CCAGGCCGACCCGGCAGCCCCGGAGCTGTGACTTCAAGGAGAACGGGgtgaggctgg	720
P R P T R Q P P E L C D F K E N G	
gggctggggggcgcgtggcgatgttcccaaggagctgaaacaggagccgtctgtgggaag	780
atgtccaggccctgggtgaggctggagctcatggatggaggagggggggtcccagtt	840
^{t3}	
gaccttgagtctcccttcagCGGGTGAAACAGTGTGAGGGACAGTCACCTGGATCA	900
R Y K Q C V G T V T L D Q	
GATCAAGGACCCGCTCGACATCACCTGCAATGAGgtgagtggcccttattgtgtcaag	960
I K D P L D I T C N E	
ttgctaattgggttgtggaaacctccctggagttaccgcgtccccatccaggc	1020
gtggaaaggccctcctacccggccctccacctcgccccaggcgtccaggctgg	1080
ctctgtcatccttagggccgcgttccctcaatgggtccccccctcgatattgtcagaa	1140
^{g3,5}	
aggcacatttcaggccccaccccgaccctctgaatcacactttgggtggagccagc	1200
tgtctttctccaagatcccagcgggttctccgtgtctgtcgctgagaggcagtac	1260
cggactaatggacttgcaggcccgtctccgtggccagcttgcgggtgggttggacc	1320
ctggcaaggccccagccatctggccctgagtcacttatgtctgtggggattcaa	1381
^{g3,5}	
ccacgtgtccaaaggcacagccagagggtggaccaggccccagccctttactgtttc	1440

FIG. 8a

cccattcaggatttctagtcggaggagggttcttgtcttgacccttgccagacc	1500
ccacccgaaacctgtttctttggcacagGTTCAAGGTGTCAGGGGAGGTCGCCGTGC	1560
F Q G V R G G R L C	
<u>G3</u>	
TATTGTAGGCCGTAGGTTCTGCGTCTGTGTCGGACGAGGATGACGGTTGCGACGGCAGGCT	1620
Y C R R F C V C V G R G ***	
<u>C5</u> <u>T5</u>	
<u>P5</u>	
TTCCCTCCCCAATTTCCGGGCCAGGTTCCGTCCCCAATTTCCGCCTCACCT	1680
TTCCGGCCCGCACCATCGGTCCACCAAGGTTCCGTAGACGGTAAGGATTGCAGG	1740
C3,5	
CAACTCACCCAGAAGGCCTTCGGTACATTAAAATCCAGCAAGGAGACCTAACGATCTG	1800
CTTGCCCAGGCCGCATCTGTCAA <u>ATAAA</u> ATTCTGTGAAACC	1843

FIG. 8b

	1 2 3	4	5 6 7 8 9	10 11 12	13	14	15 16	17 18
PG-1	RGG	R	LCYCR	RRF	C	V	CV	GR*
PG-2	RGG	R	LCYCR	RRF	C	I	CV	GR*
PG-3	RGG	G	LCYCR	RRF	C	V	CV	GR*
PG-4	RGG	R	LCYCR	GWI	C	F	CV	GR*
PG-5	RGG	R	LCYCR	PRF	C	V	CV	GR*

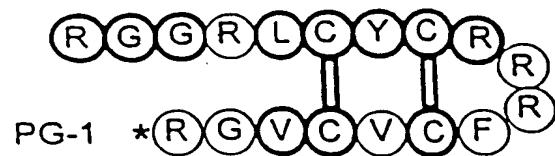


FIG. 10

FIG. II a-2

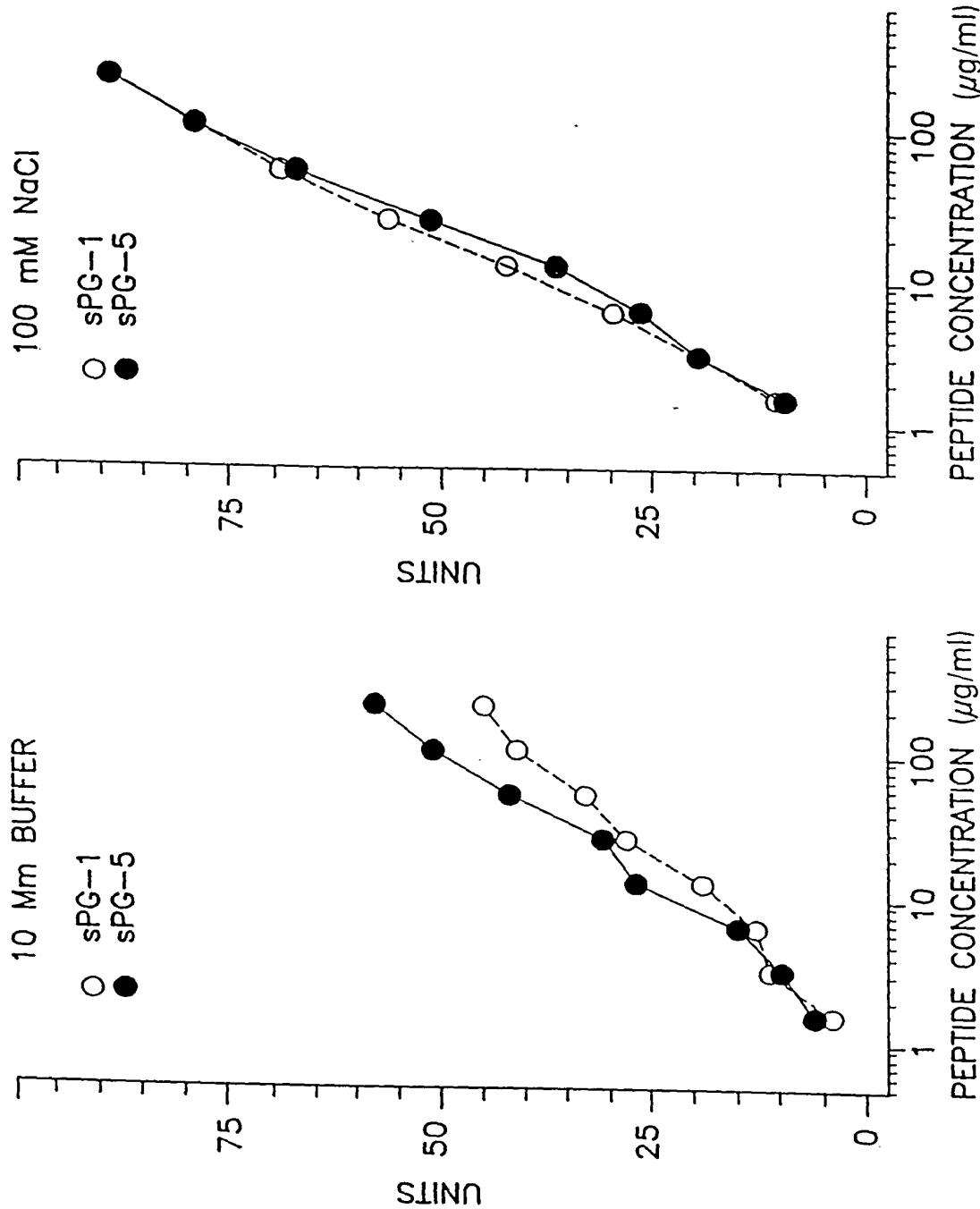


FIG. II a-1

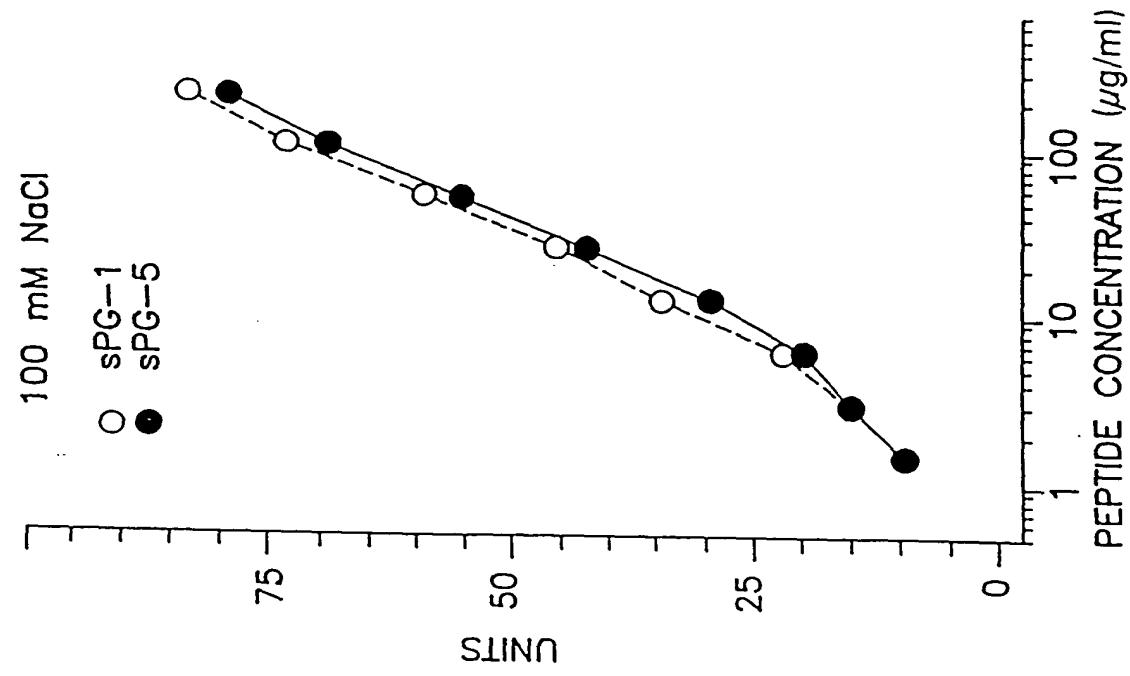
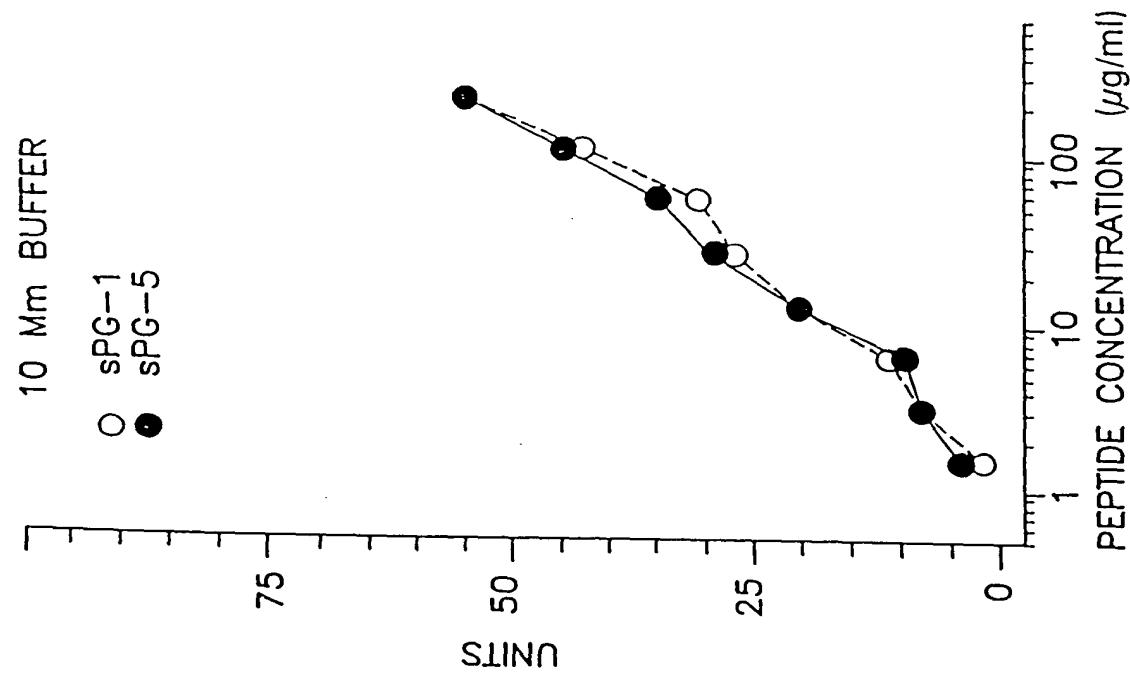


FIG. IIb-1

FIG. IIb-2

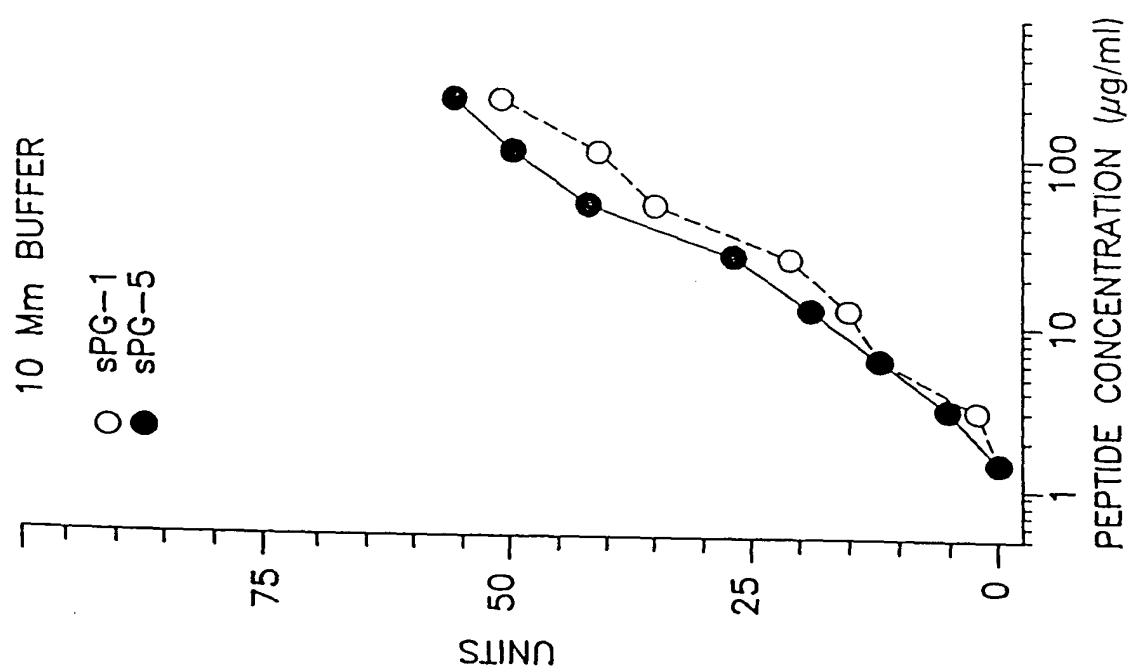


FIG. II c-1

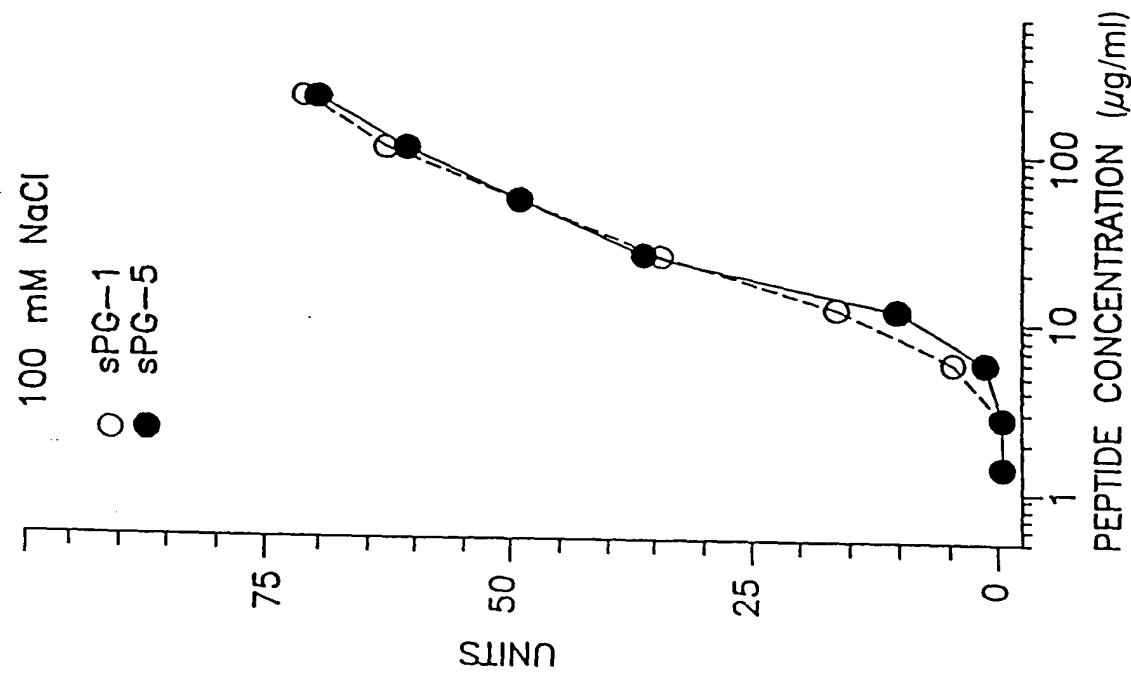


FIG. II c-2

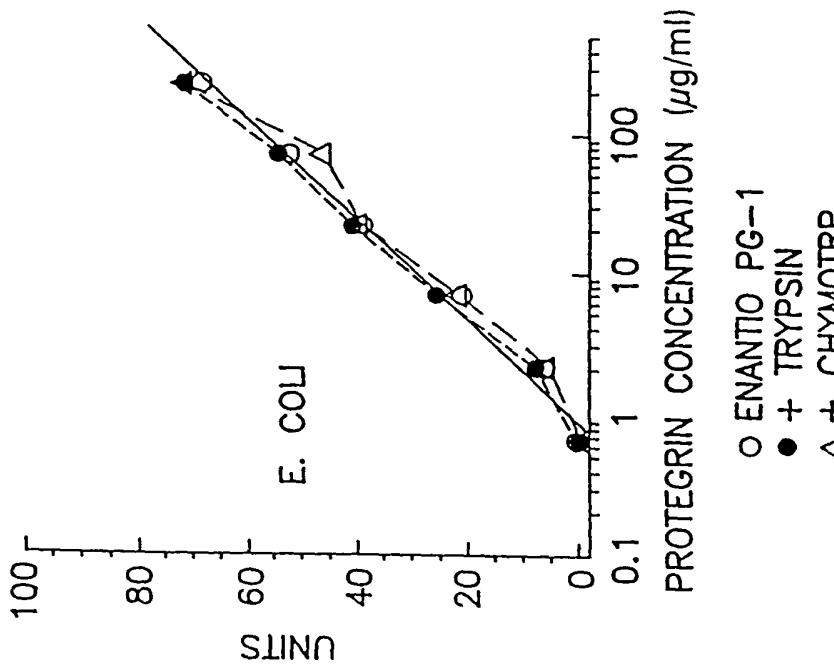


FIG. 12b

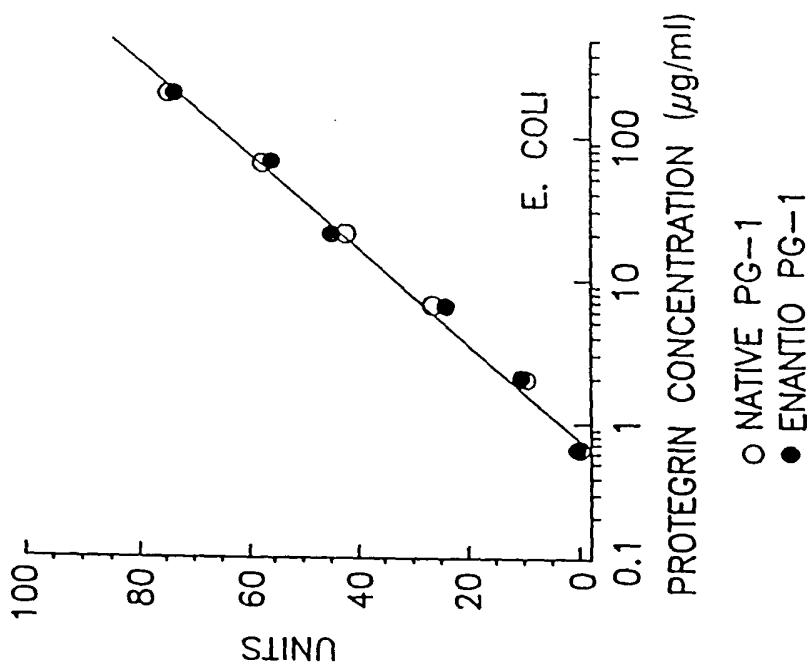


FIG. 12a

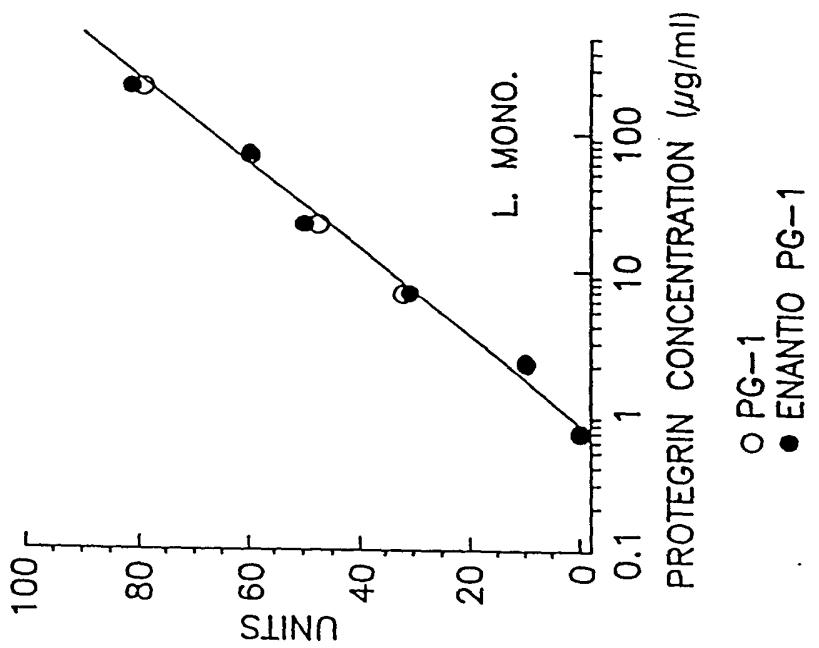


FIG. 12d

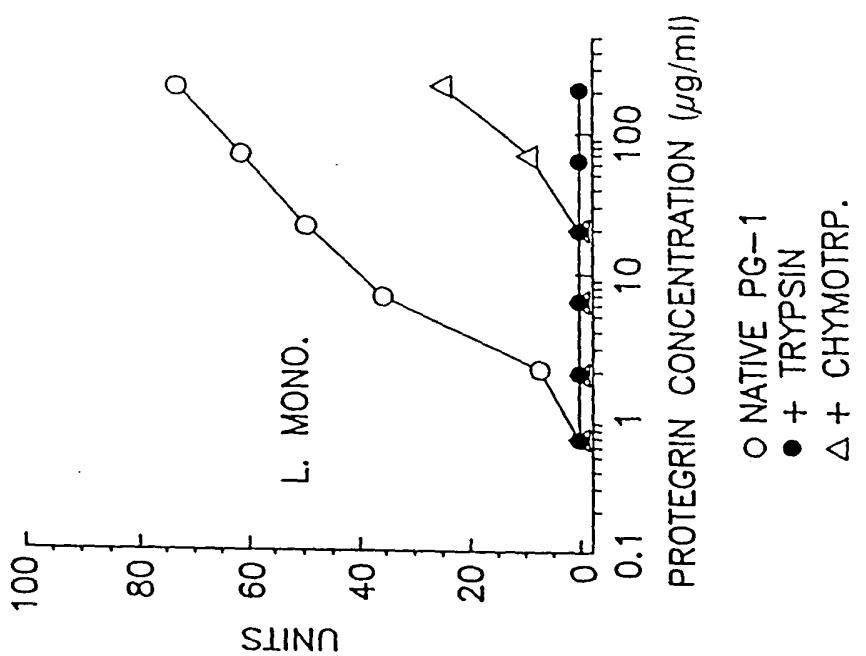


FIG. 12c

OPEN SYMBOLS = KITE, CLOSED SYMBOLS = BULLET

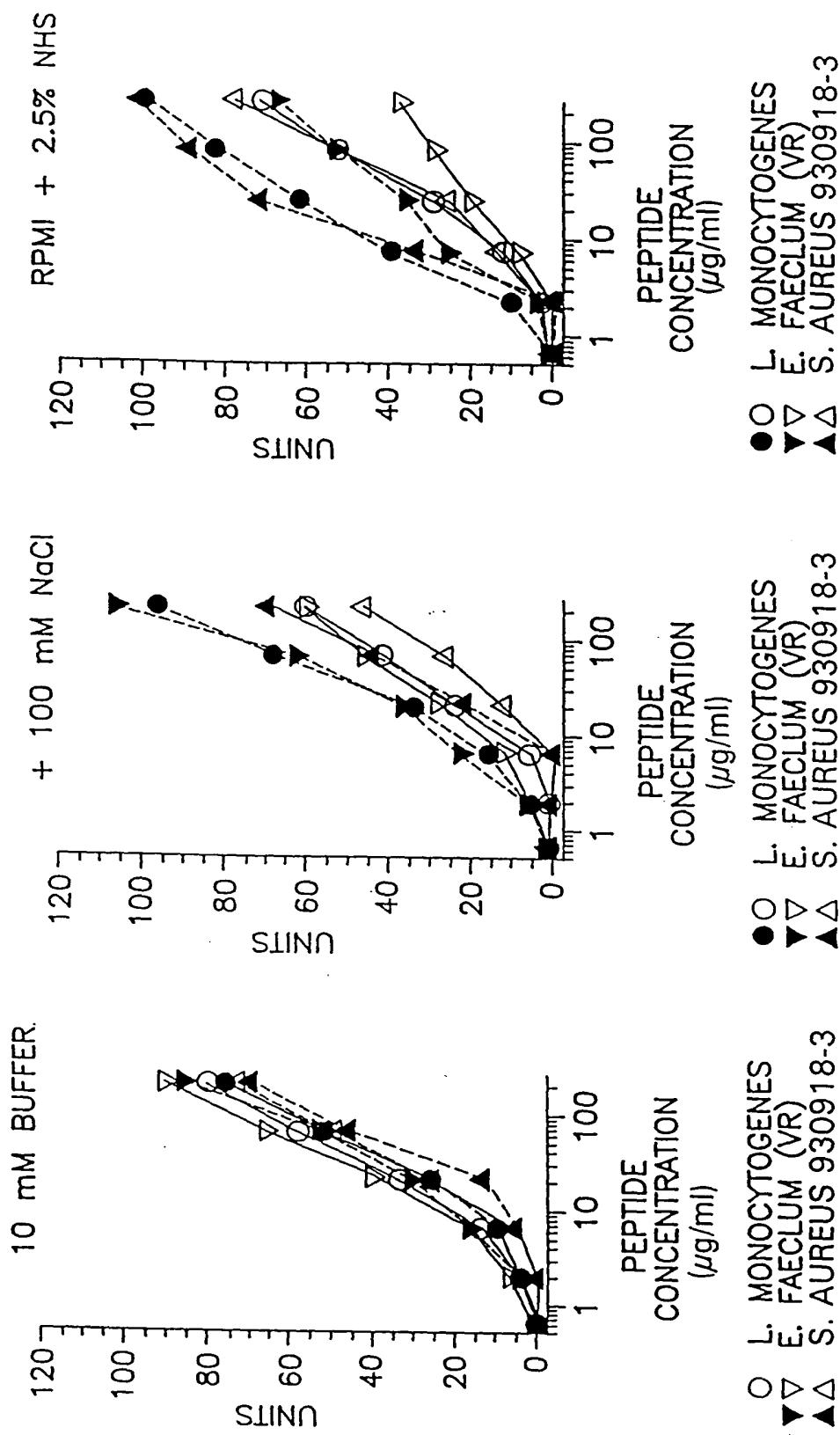


FIG. I3a

FIG. I3b

FIG. I3c

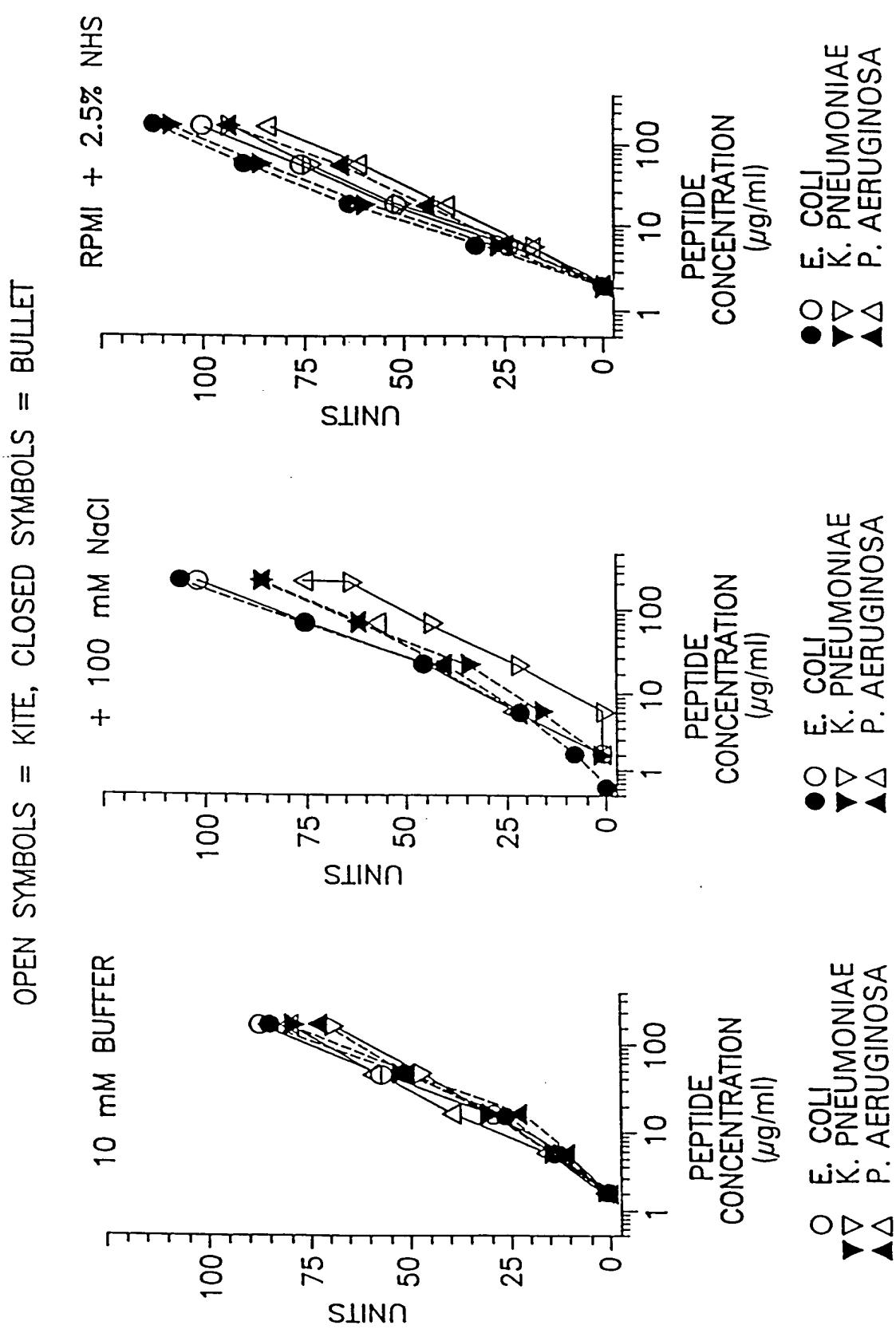


FIG. 14a

FIG. 14b

FIG. 14c

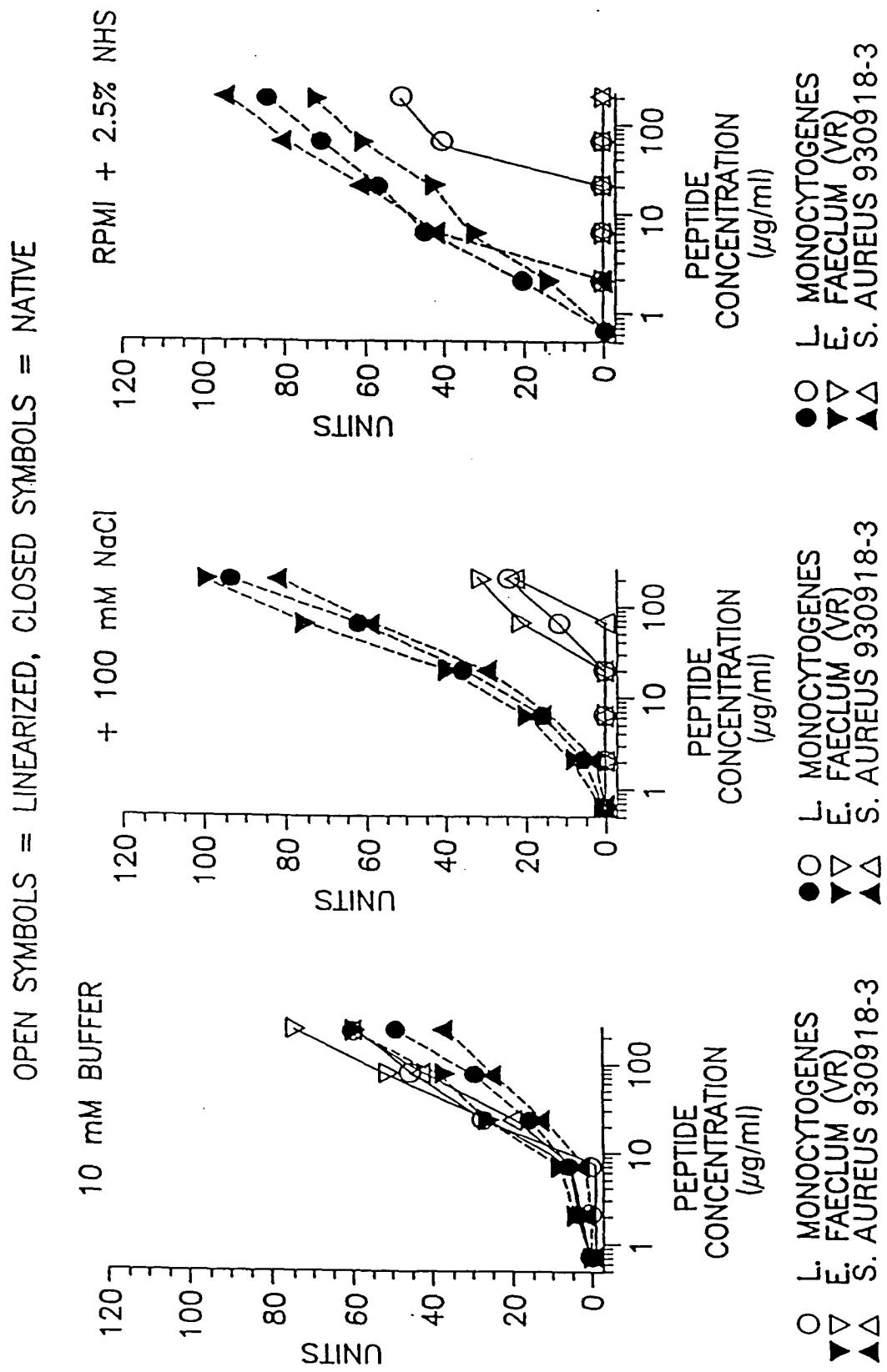


FIG. 15a

FIG. 15b

FIG. 15c

OPEN SYMBOLS = LINEARIZED, CLOSED SYMBOLS = sPG-1

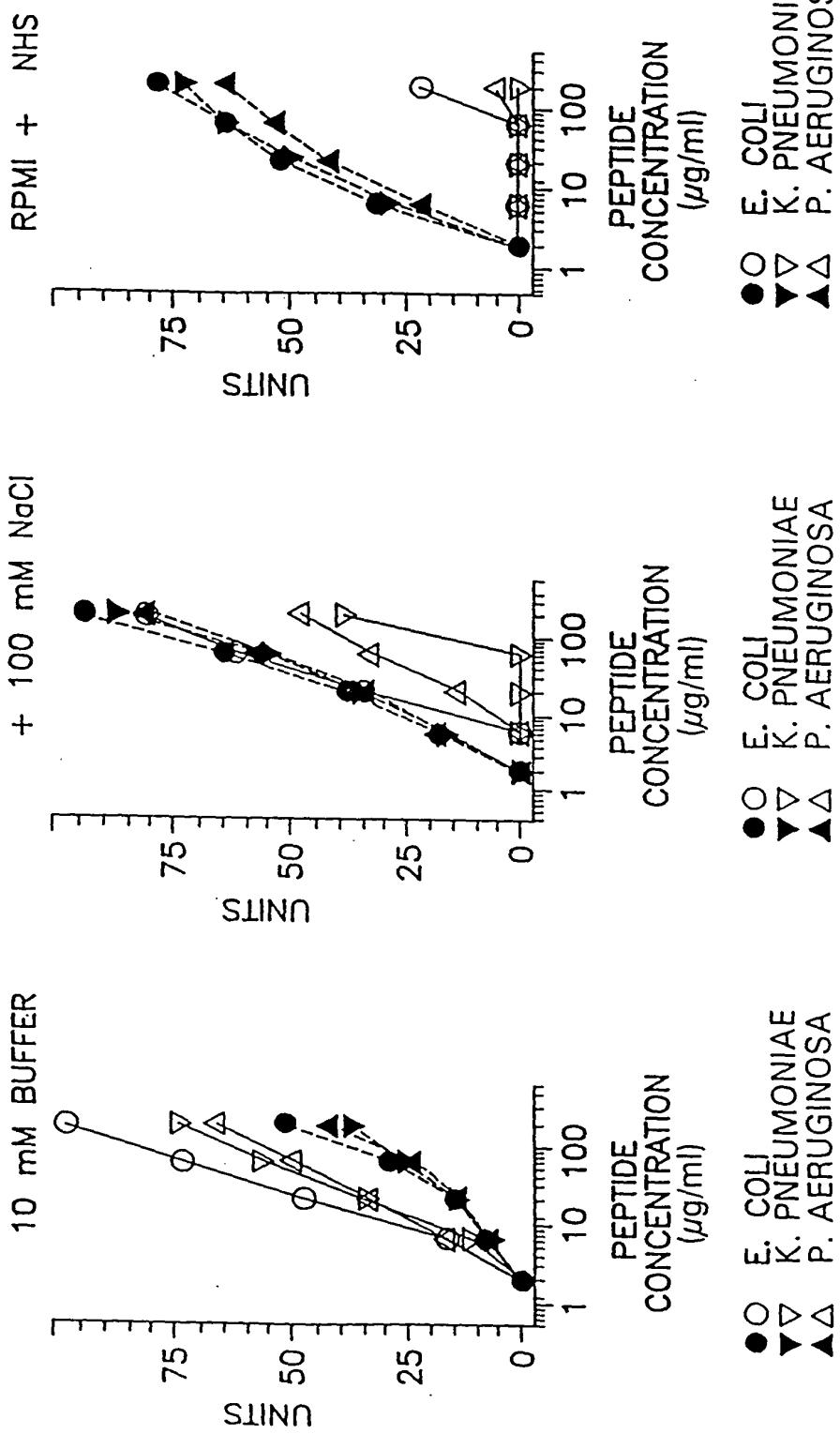


FIG. 16a

FIG. 16b

FIG. 16c